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SEQUENCE LISTING

<110> Olsen, Carsten
Rasmussen, Michael Dolberg

<120> Method for Stable Gene-Amplification in a Bacterial Host Cell

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 Phe Gly Arg His Asp Val Asp Leu Pro Trp Glu Arg Thr Asp Lys Ala
 375 380 385 390
 gag cag ctg cgt aaa gaa gcg tta gga gaa taa ttttatagcc gcttactggt 2219
 Glu Gln Leu Arg Lys Glu Ala Leu Gly Glu
 395 400
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Ile Leu Lys Lys Asp Pro Asn Ala Arg Val Ala Cys Glu Thr Ser Val
 35 40 45

Thr Thr Gly Leu Val Leu Val Ser Gly Glu Ile Thr Thr Ser Thr Tyr
 50 55 60

Val Asp Ile Pro Lys Thr Val Arg Gln Thr Ile Lys Glu Ile Gly Tyr
 65 70 75 80

Thr Arg Ala Lys Tyr Gly Phe Asp Ala Glu Thr Cys Ala Val Leu Thr
 85 90 95

Ser Ile Asp Glu Gln Ser Ala Asp Ile Ala Met Gly Val Asp Pro Ala
100 105 110

Leu Glu Ala Arg Glu Gly Thr Met Ser Asp Glu Glu Ile Glu Ala Ile
115 120 125

Gly Ala Gly Asp Gln Gly Leu Met Phe Gly Tyr Val Cys Asn Glu Thr
130 135 140

Lys Glu Leu Met Pro Leu Pro Ile Ser Leu Ala His Lys Leu Ala Arg
145 150 155 160

Arg Leu Ser Glu Val Arg Lys Glu Asp Ile Leu Pro Tyr Leu Arg Pro
165 170 175

Asp Gly Lys Thr Gln Val Thr Val Glu Tyr Asp Glu Asn Asn Lys Pro
180 185 190

Val Arg Ile Asp Ala Ile Val Ile Ser Thr Gln His His Pro Glu Ile
195 200 205

Thr Leu Glu Gln Ile Gln Arg Asn Ile Lys Glu His Val Ile Asn Pro
210 215 220

Val Val Pro Glu Glu Leu Ile Asp Glu Glu Thr Lys Tyr Phe Ile Asn
225 230 235 240

Pro Thr Gly Arg Phe Val Ile Gly Gly Pro Gln Gly Asp Ala Gly Leu
245 250 255

Thr Gly Arg Lys Ile Ile Val Asp Thr Tyr Gly Gly Tyr Ala Arg His
260 265 270

Gly Gly Gly Ala Phe Ser Gly Lys Asp Ala Thr Lys Val Asp Arg Ser
275 280 285

Ala Ala Tyr Ala Ala Arg Tyr Val Ala Lys Asn Ile Val Ala Ala Glu
290 295 300

Leu Ala Asp Ser Cys Glu Val Gln Leu Ala Tyr Ala Ile Gly Val Ala
305 310 315 320

Gln Pro Val Ser Ile Ser Ile Asn Thr Phe Gly Ser Gly Lys Ala Ser
 325 330 335

Glu Glu Lys Leu Ile Glu Val Val Arg Asn Asn Phe Asp Leu Arg Pro
 340 345 350

Ala Gly Ile Ile Lys Met Leu Asp Leu Arg Arg Pro Ile Tyr Lys Gln
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Thr Ala Ala Tyr Gly His Phe Gly Arg His Asp Val Asp Leu Pro Trp
 370 375 380

Glu Arg Thr Asp Lys Ala Glu Gln Leu Arg Lys Glu Ala Leu Gly Glu
 385 390 395 400